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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Tue Jul 17 13:54:26 EDT 2007

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Reviewer Comments:

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11

The above <222> (4)..(4) and the <223> response under it are
incorrect: "c," not "n," is at location 4, and can only represent itself.

Application No: 10593357

Version No: 1.1

Input Set:**Output Set:****Started:** 2007-07-16 13:54:05.493**Finished:** 2007-07-16 13:54:08.628**Elapsed:** 0 hr(s) 0 min(s) 3 sec(s) 135 ms**Total Warnings:** 86**Total Errors:** 0**No. of SeqIDs Defined:** 86**Actual SeqID Count:** 86

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Input Set:

Output Set:

Started: 2007-07-16 13:54:05.493
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Total Errors: 0
No. of SeqIDs Defined: 86
Actual SeqID Count: 86

Error code

Error Description

This error has occurred more than 20 times, will not be displayed

SEQUENCE LISTING

<110> BRAIN BIOTECHNOLOGY RESEARCH AND INFORMATION NETWORK AG
 Verseck, Stefan
 Liebeton, Klaus
 Eck, Jorgen

<120> Nitrile hydratases from metagenome libraries

<130> 009848-0356700

<140> 10/593,357

<141> 2006-09-18

<150> WO PCT/EP2005/002556

<151> 2005-03-10

<150> DE 10 2004 013 842.7

<151> 2004-03-20

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48

Met Ser Lys His Val His Asp Tyr Tyr Ala Lys Lys Lys His Asp His

1

5

10

15

gat cat gac cac gac gtc cac gag gcg atc gag gac cgg gac gag ggt	96
Asp His Asp His Asp Val His Glu Ala Ile Glu Asp Arg Asp Glu Gly	
20 25 30	
ccg ccg tcg gaa ttc gag atc atg agc cgc gcc atg cag gag ctg ctg	144
Pro Pro Ser Glu Phe Glu Ile Met Ser Arg Ala Met Gln Glu Leu Leu	
35 40 45	
gaa gag aag ggc gtc gtc acc gcg gag cag gtc cgg cgc agc atg gag	192
Glu Glu Lys Gly Val Val Thr Ala Glu Gln Val Arg Arg Ser Met Glu	
50 55 60	
aag ttc gag gaa gag ctg ccc tac cgg ggg gcg cgg gtc gtc gcg cac	240
Lys Phe Glu Glu Glu Leu Pro Tyr Arg Gly Ala Arg Val Val Ala His	
65 70 75 80	
gcc tgg acc gac ccg gaa ttc aag aag cgg ctg ctg gcg gac ggc aag	288
Ala Trp Thr Asp Pro Glu Phe Lys Lys Arg Leu Leu Ala Asp Gly Lys	
85 90 95	
gcc gcc gtc tcg gag ttc ggc atc gat ttg gaa gcc gag cgg ctg atc	336
Ala Ala Val Ser Glu Phe Gly Ile Asp Leu Glu Ala Glu Arg Leu Ile	
100 105 110	
gcg gtc gcg aac aca acg gac gtg cac aac gtc atc gtg tgc acg ctg	384
Ala Val Ala Asn Thr Thr Asp Val His Asn Val Ile Val Cys Thr Leu	
115 120 125	
tgc tcg tgc tac ccg cgc acg ctg ctc ggc atg ccg ccg acc tgg tac	432
Cys Ser Cys Tyr Pro Arg Thr Leu Leu Gly Met Pro Pro Thr Trp Tyr	
130 135 140	
aag agc gat aac tac cgc tcg cgc gtg gtc tac gaa ccg cgc gcg gtg	480
Lys Ser Asp Asn Tyr Arg Ser Arg Val Val Tyr Glu Pro Arg Ala Val	
145 150 155 160	
ctg aag gaa ttc ggc acc gtg ctg ccg gag cgc gtc acc gtg cgc gtg	528
Leu Lys Glu Phe Gly Thr Val Leu Pro Glu Arg Val Thr Val Arg Val	
165 170 175	
cac gac tcc aac gcc gac atg cgc tac gtg gtg atc ccc atg cgc ccg	576
His Asp Ser Asn Ala Asp Met Arg Tyr Val Val Ile Pro Met Arg Pro	
180 185 190	
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Gln Gly Thr Glu Gly Trp Ser Glu Glu Arg Leu Ala Glu Leu Leu Thr	
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cgc gac acg ctg gtg ggg gtc acc gtg cca aaa gtg gaa gtc gga agt	672
Arg Asp Thr Leu Val Gly Val Thr Val Pro Lys Val Glu Val Gly Ser	
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Pro Pro Ser Glu Phe Glu Ile Met Ser Arg Ala Met Gln Glu Leu Leu
35 40 45

Glu Glu Lys Gly Val Val Thr Ala Glu Gln Val Arg Arg Ser Met Glu
50 55 60

Lys Phe Glu Glu Glu Leu Pro Tyr Arg Gly Ala Arg Val Val Ala His
65 70 75 80

Ala Trp Thr Asp Pro Glu Phe Lys Lys Arg Leu Leu Ala Asp Gly Lys
85 90 95

Ala Ala Val Ser Glu Phe Gly Ile Asp Leu Glu Ala Glu Arg Leu Ile
100 105 110

Ala Val Ala Asn Thr Thr Asp Val His Asn Val Ile Val Cys Thr Leu
115 120 125

Cys Ser Cys Tyr Pro Arg Thr Leu Leu Gly Met Pro Pro Thr Trp Tyr
130 135 140

Lys Ser Asp Asn Tyr Arg Ser Arg Val Val Tyr Glu Pro Arg Ala Val
145 150 155 160

Leu Lys Glu Phe Gly Thr Val Leu Pro Glu Arg Val Thr Val Arg Val
165 170 175

His Asp Ser Asn Ala Asp Met Arg Tyr Val Val Ile Pro Met Arg Pro
180 185 190

Gln Gly Thr Glu Gly Trp Ser Glu Glu Arg Leu Ala Glu Leu Leu Thr
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Arg Lys Ser Lys Gly Gly Ser Lys Thr Arg
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 Ser Ala Met Asn Ile Pro Ala Arg Glu Phe Ala Leu Gln Arg Thr Ala
 20 25 30

ccg gtc gag cag cgt gtc gac gcg atc cag gcg gcg ctc gac gaa cgc 144
 Pro Val Glu Gln Arg Val Asp Ala Ile Gln Ala Ala Leu Asp Glu Arg
 35 40 45

ggc ttg aac gcc agt gac gca gtc cag gaa ttg agc cac ctg gcg gag 192
 Gly Leu Asn Ala Ser Asp Ala Val Gln Glu Leu Ser His Leu Ala Glu
 50 55 60

gag caa tgg att ccg cgc aat ggc gcg cgg gtc gtc gcc aaa gcc tgg 240
 Glu Gln Trp Ile Pro Arg Asn Gly Ala Arg Val Val Ala Lys Ala Trp
 65 70 75 80

gtc gac ccg gaa ttc cgc gcg cgg ctt ctg gcc gac ggt cgc gcc gcc 288
 Val Asp Pro Glu Phe Arg Ala Arg Leu Leu Ala Asp Gly Arg Ala Ala
 85 90 95

gtt gcc gaa ctg ggc ctc tcg atg ccg aag cat cac cgg cac ctc gtg 336
 Val Ala Glu Leu Gly Leu Ser Met Pro Lys His His Arg His Leu Val
 100 105 110

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Val Leu Glu Asn Thr Pro Ser Val Gln Asn Val Ile Cys Cys Thr Gln
115 120 125

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Cys Ser Cys Thr Ala Phe Thr Ile Ile Gly Leu Pro Pro Asp Trp Tyr
130 135 140

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Lys Asp Leu Glu Tyr Arg Ala Arg Val Val Arg Glu Ser Arg Thr Val
145 150 155 160

ctg aag gag atg gga ctg gat ctg cct cgg gat gtc gaa att cgc gtc 528
Leu Lys Glu Met Gly Leu Asp Leu Pro Arg Asp Val Glu Ile Arg Val
165 170 175

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